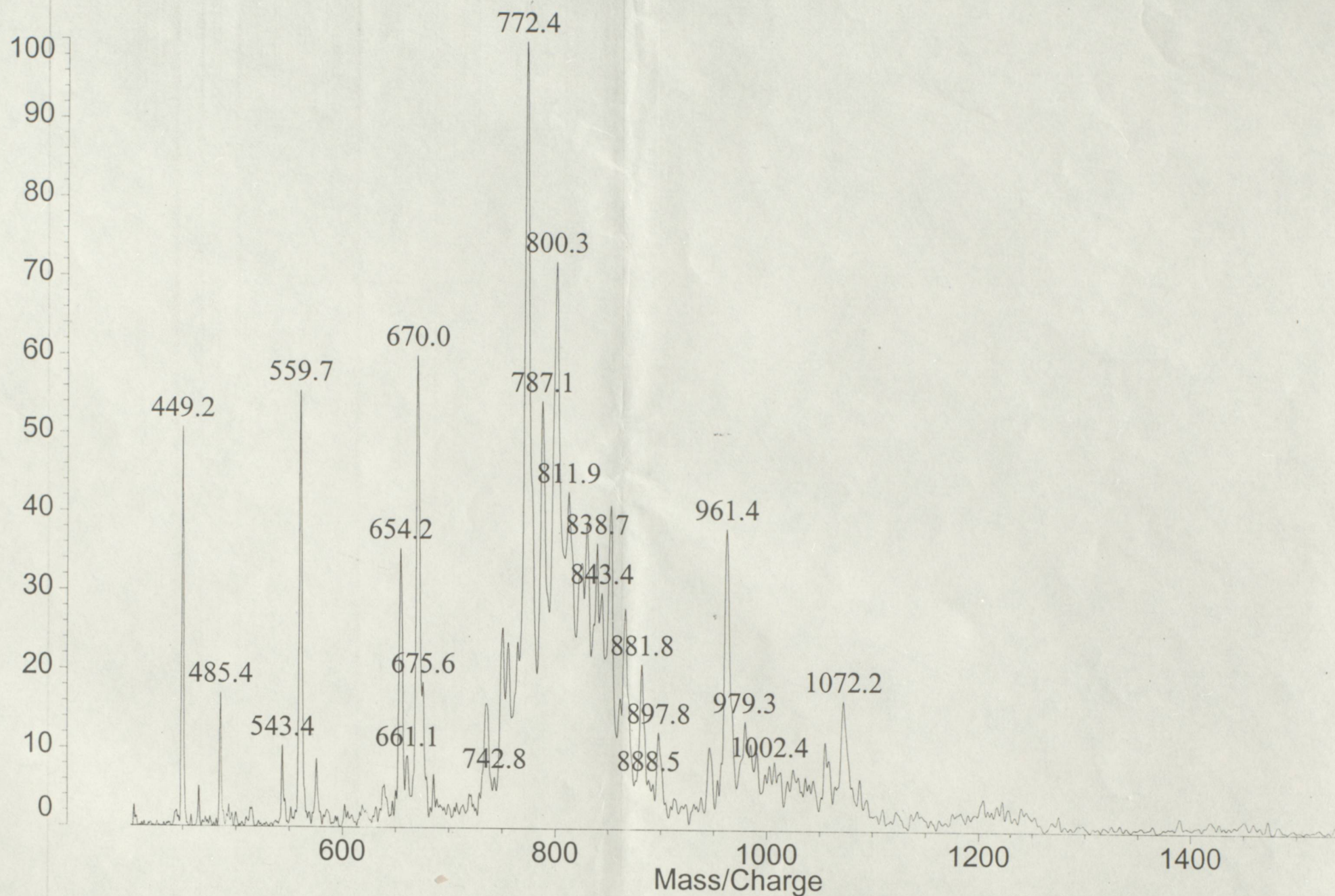
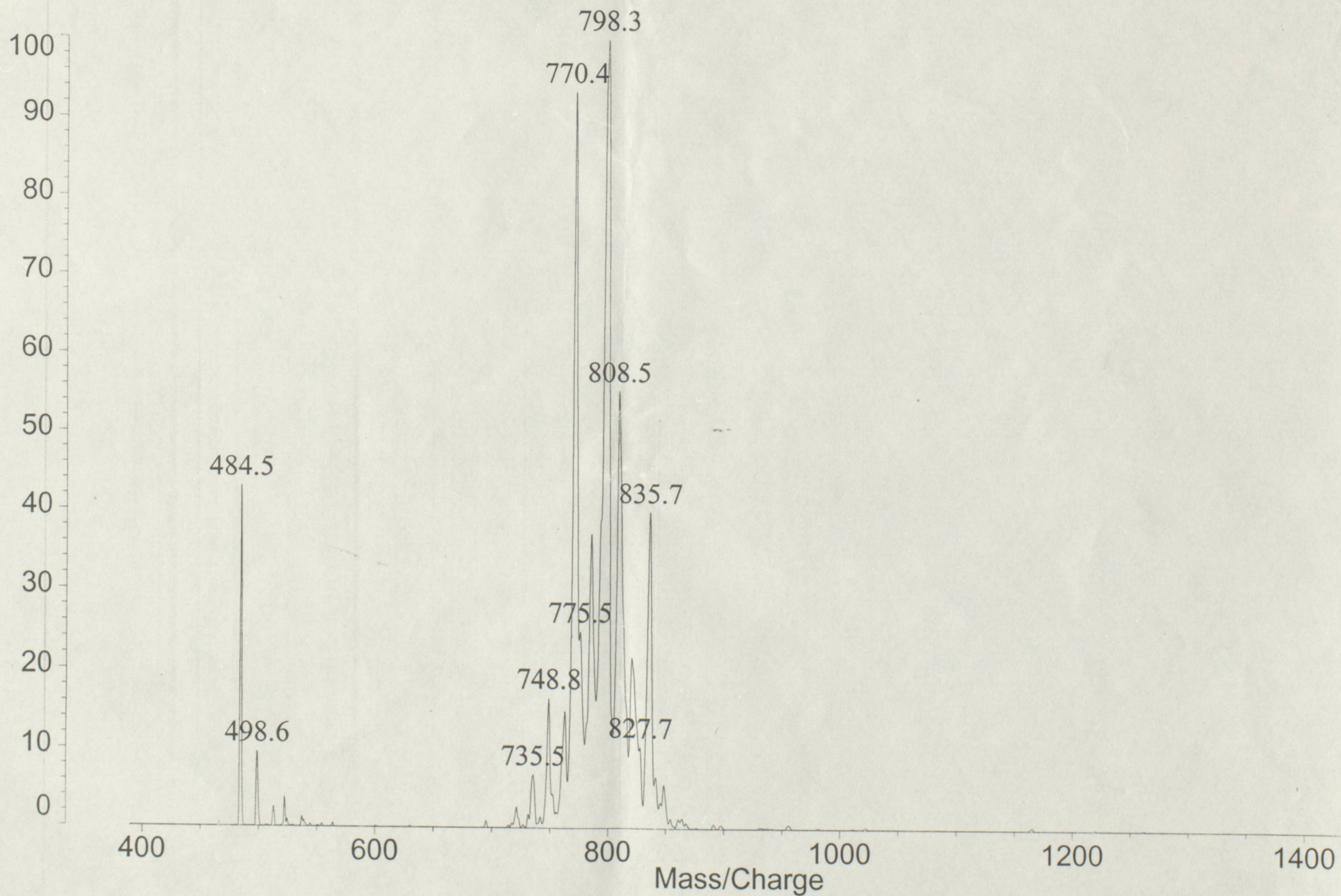


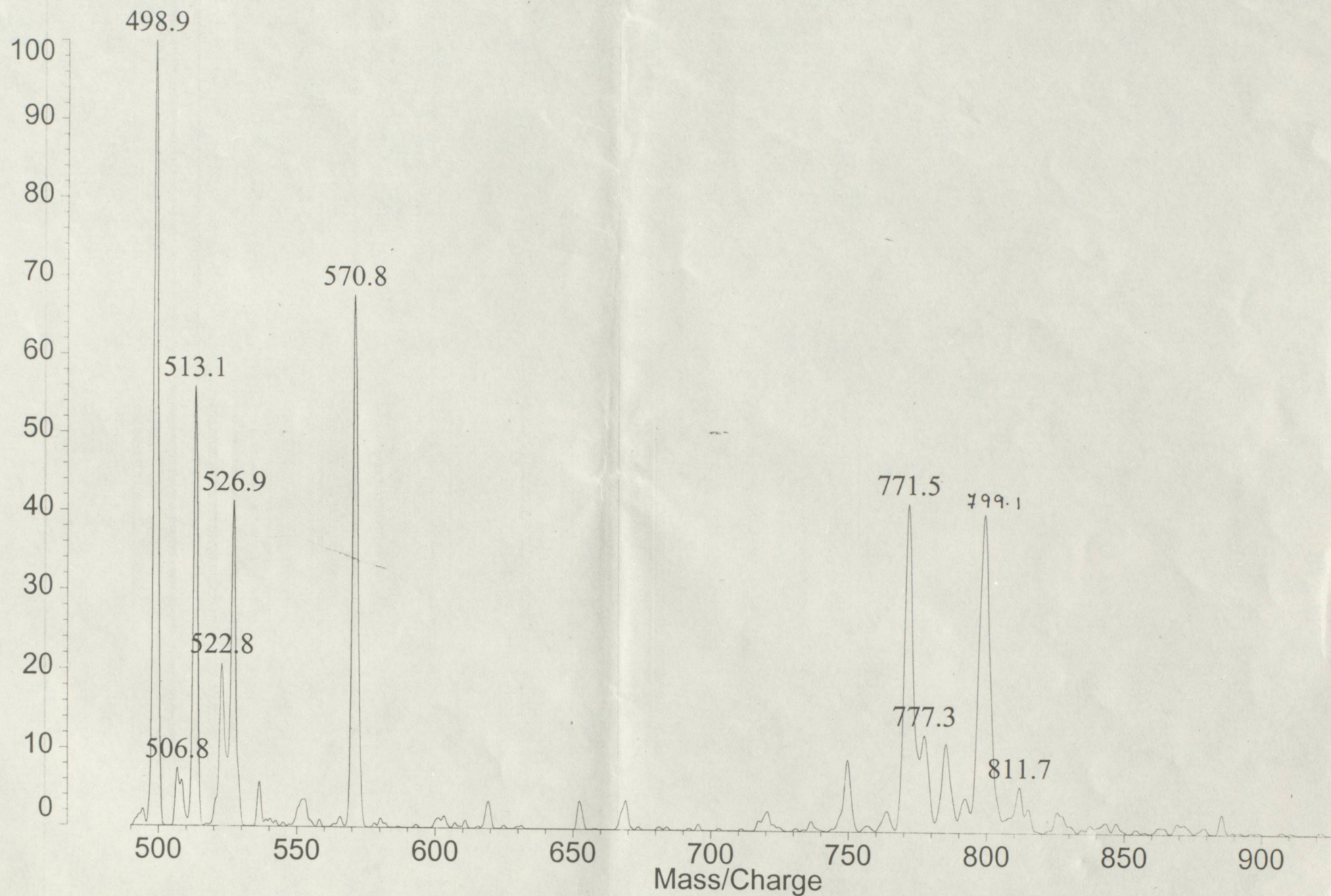
%Int. 100% = 6.9 mV[sum= 481 mV] Profiles 1-70 Smooth Av 200



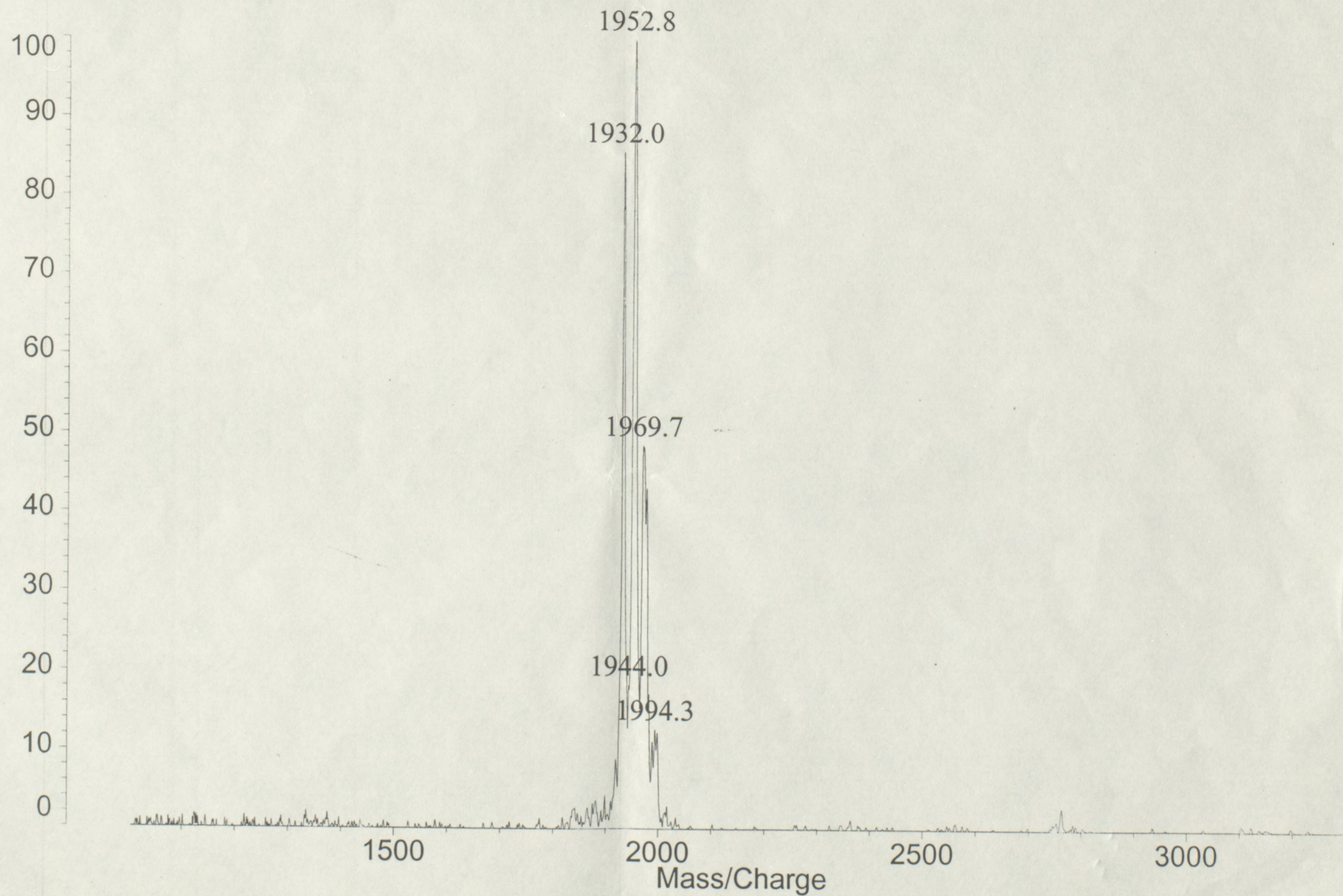
%Int. 100% = 6.3 mV[sum= 348 mV] Profiles 1-55 Smooth Av 200



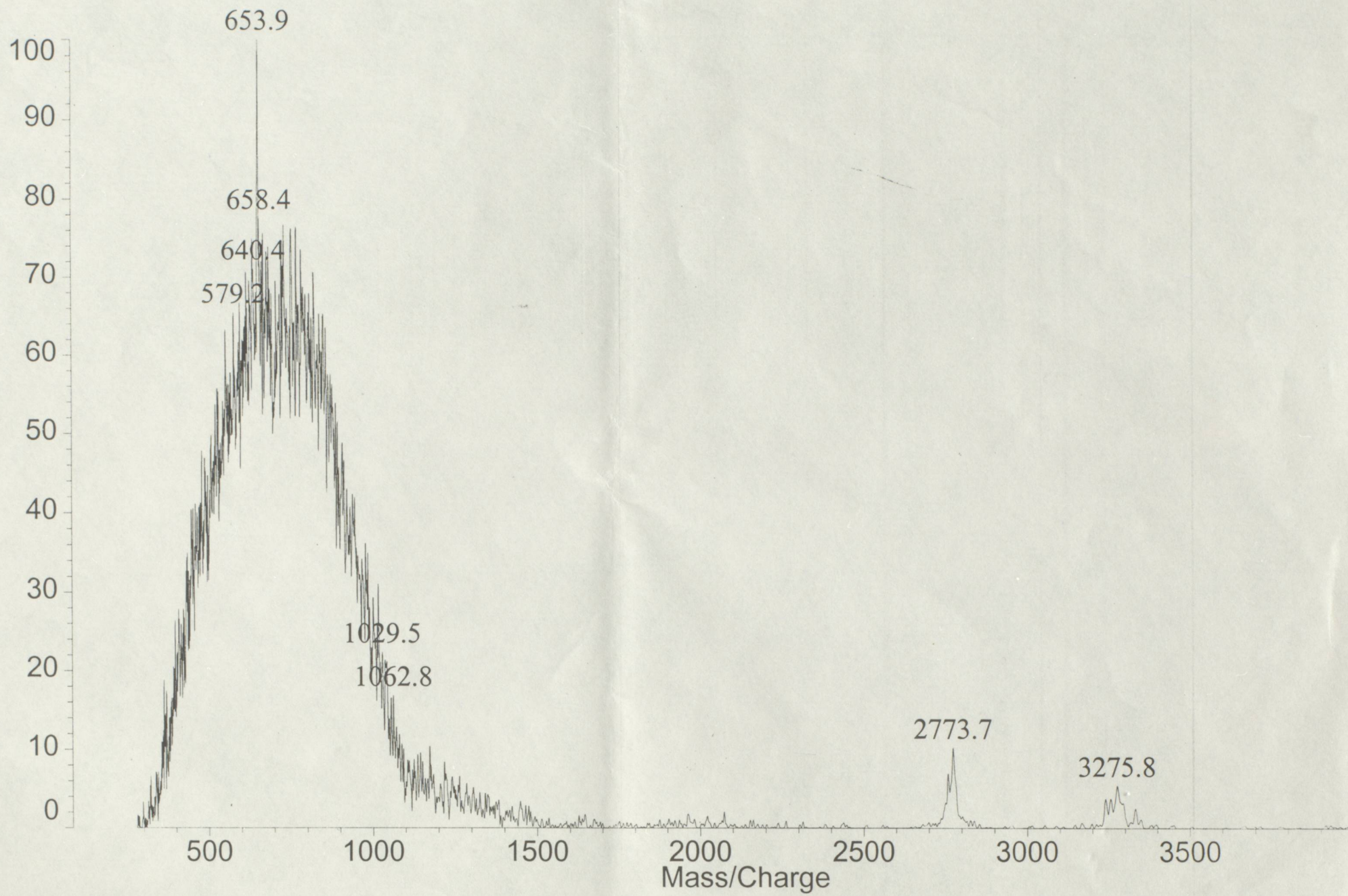
%Int. 100% = 7.0 mV[sum= 443 mV] Profiles 1-63 Smooth Av 200



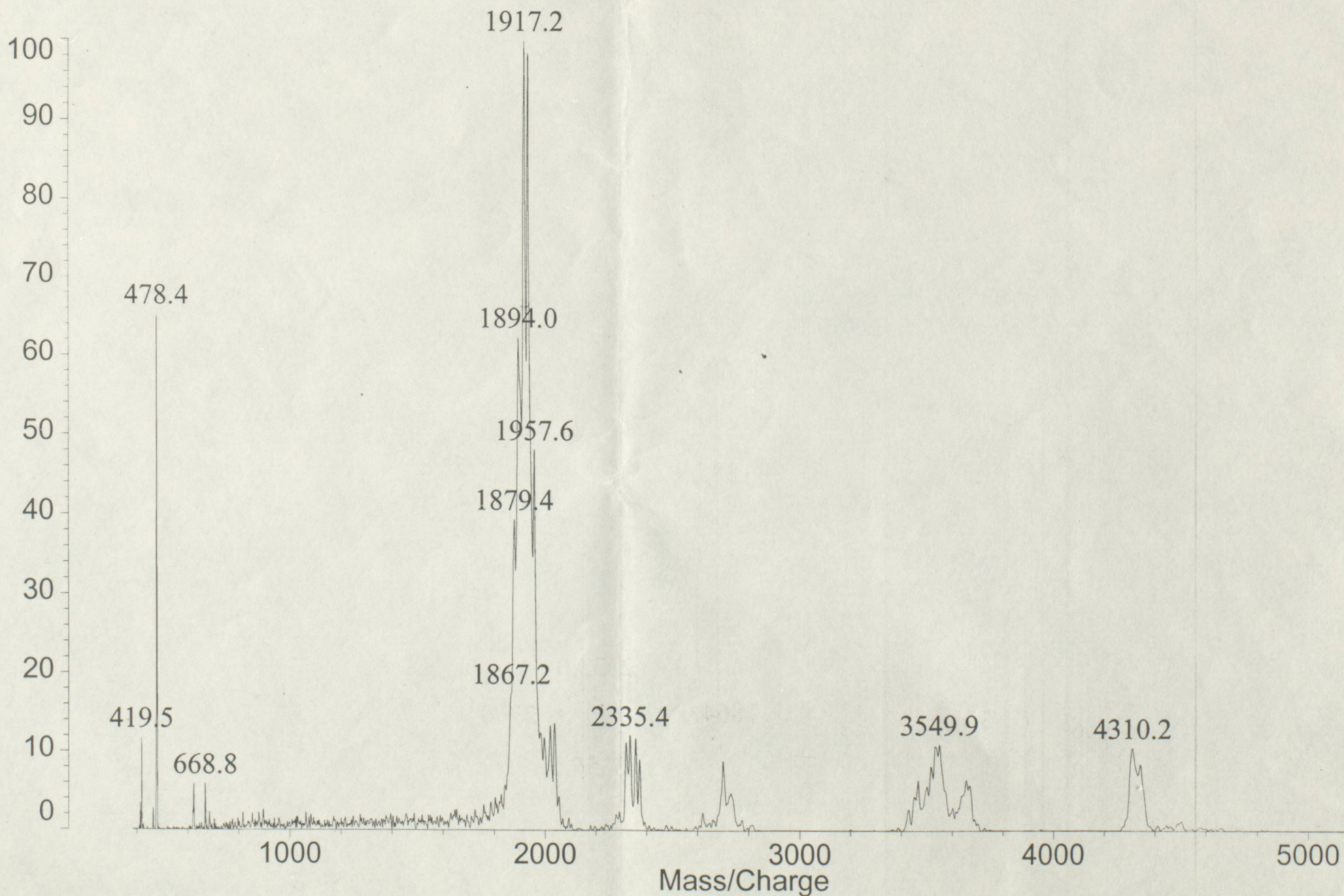
%Int. 100% = 4.0 mV[sum= 279 mV] Profiles 1-70 Smooth Av 100



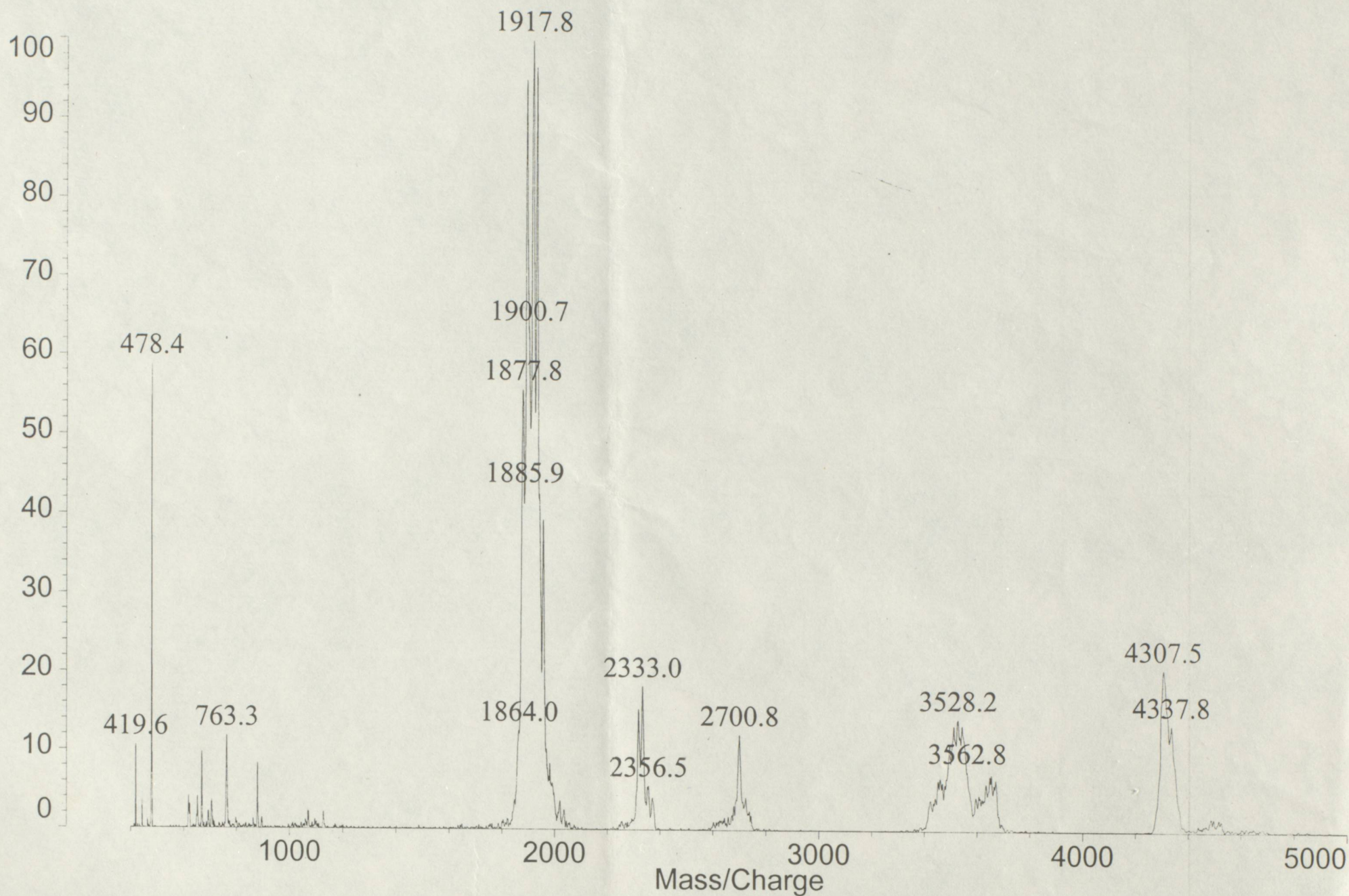
%Int. 100% = 3.4 mV[sum= 237 mV] Profiles 1-70 Smooth Av 200



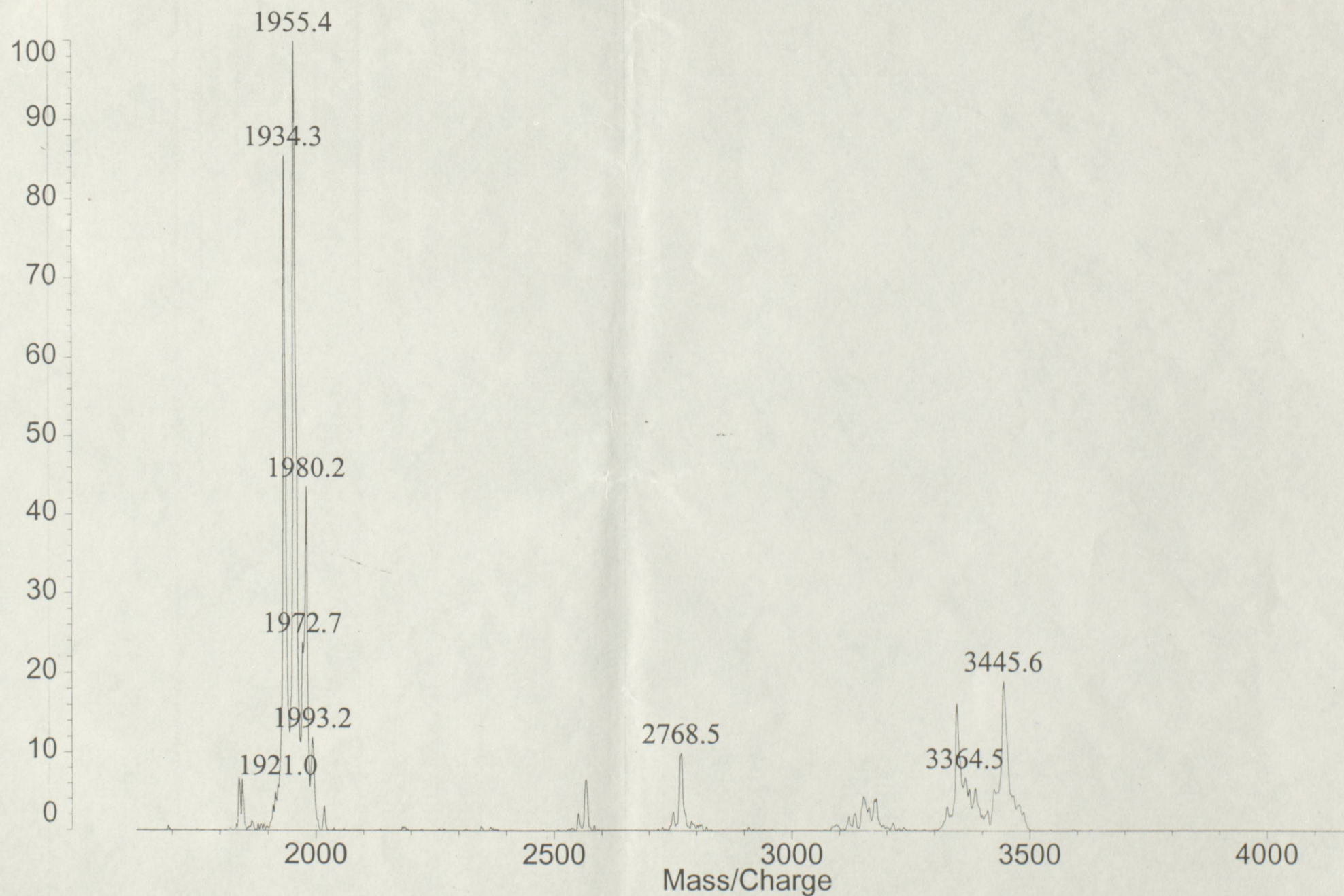
%Int. 100% = 51 mV[sum= 508 mV] Profiles 1-10 Smooth Av 200



%Int. 100% = 27 mV[sum= 1074 mV] Profiles 1-40 Smooth Av 100

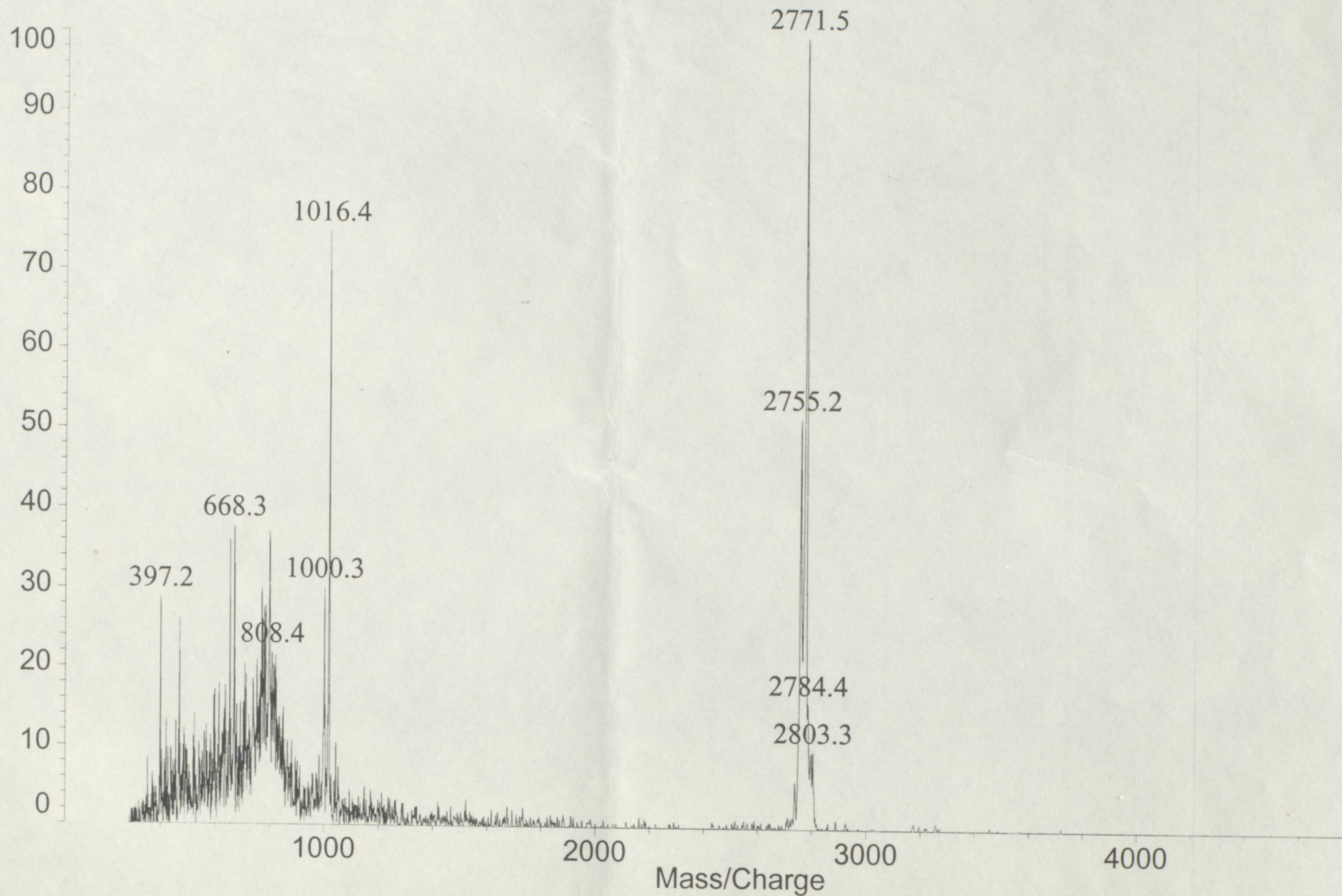


%Int. 100% = 22 mV[sum= 817 mV] Profiles 1-38 Smooth Av 100

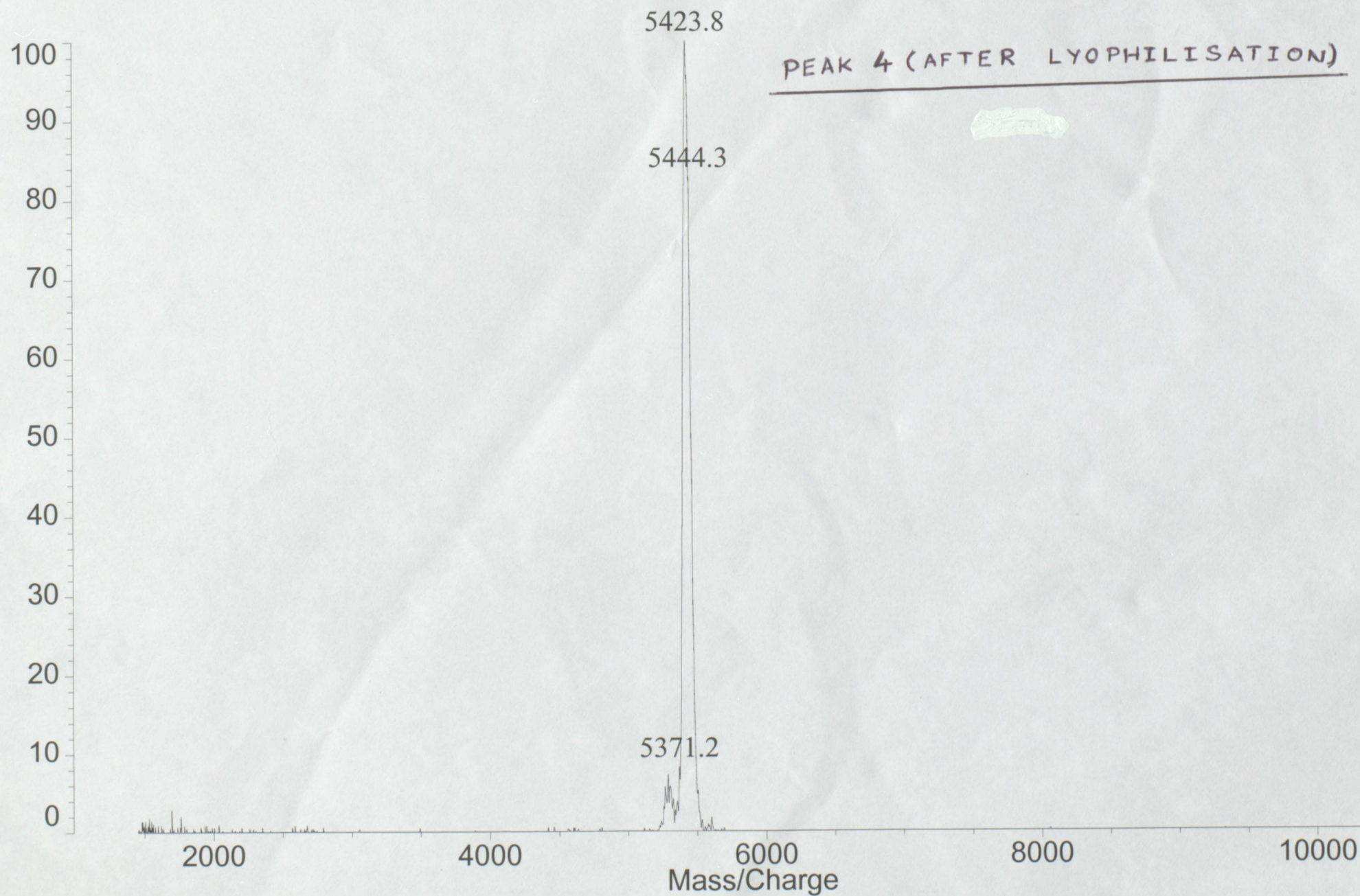


Data: <Untitled>.14 1 Jul 2001 21:24 Cal: peptide\_calibration 13 Feb 2000 10:56  
Kratos PCKompact SEQ V1.2.2: + Linear High, Power: 130, P.Ext. @ 2500 (bin 153) **BOILED** **AMADIS**  
(solution only)

%Int. 100% = 3.1 mV[sum= 217 mV] Profiles 1-70 Smooth Av 100



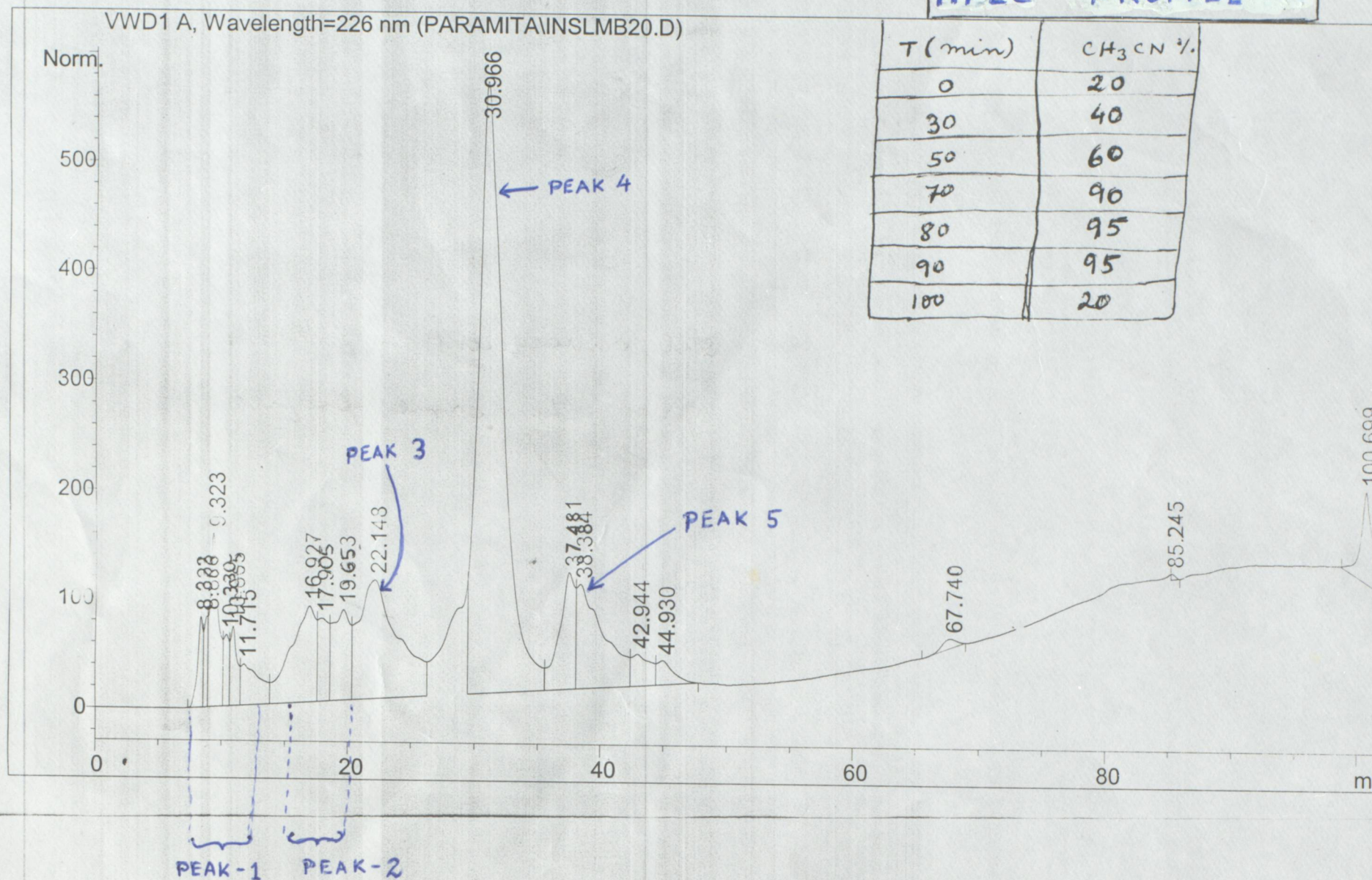
%Int. 100% = 7.5 mV[sum= 298 mV] Profiles 1-40 Smooth Av 100



16th July '01

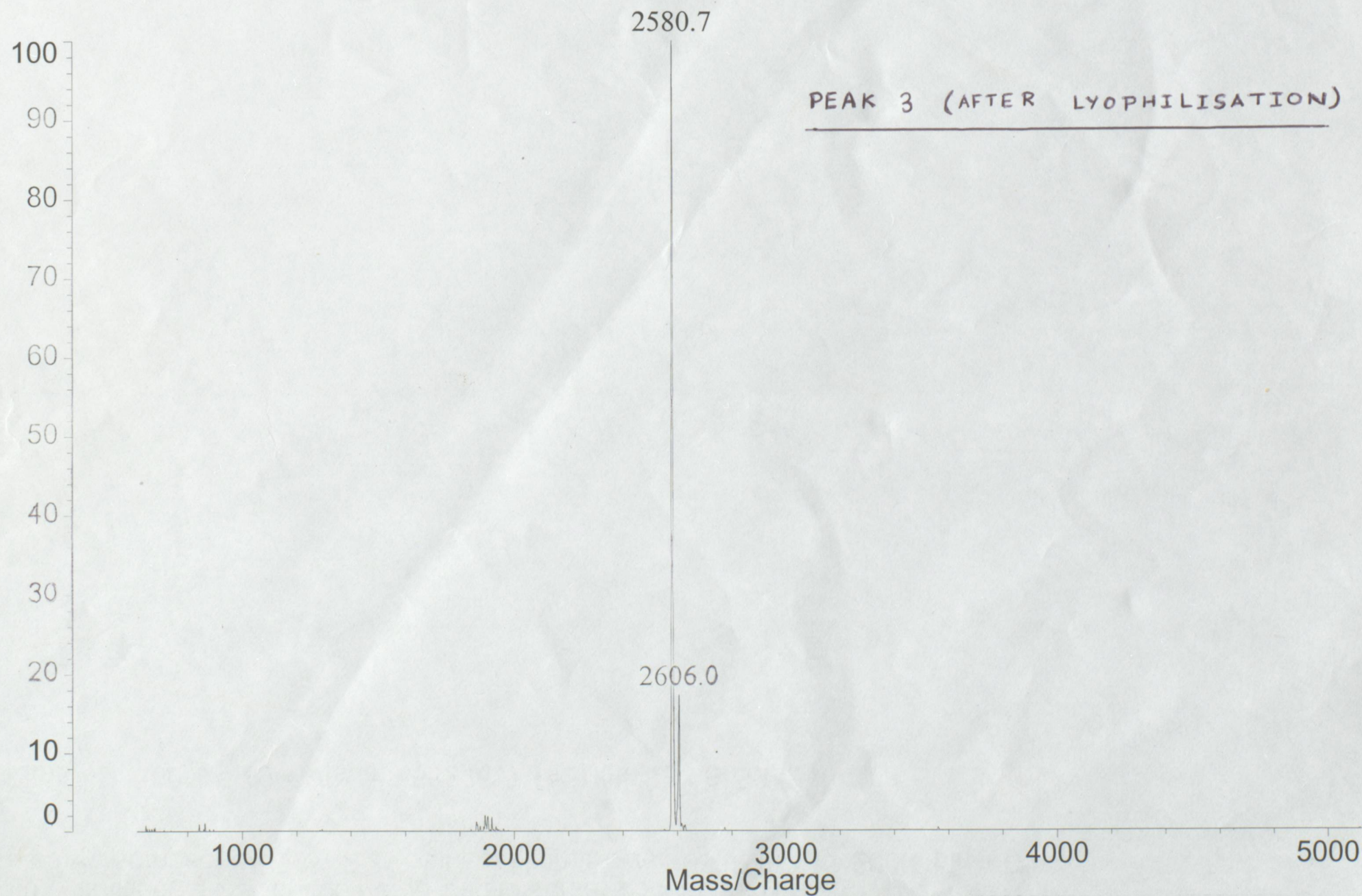
# HPLC PROFILE

VWD1 A, Wavelength=226 nm (PARAMITA\INSLMB20.D)

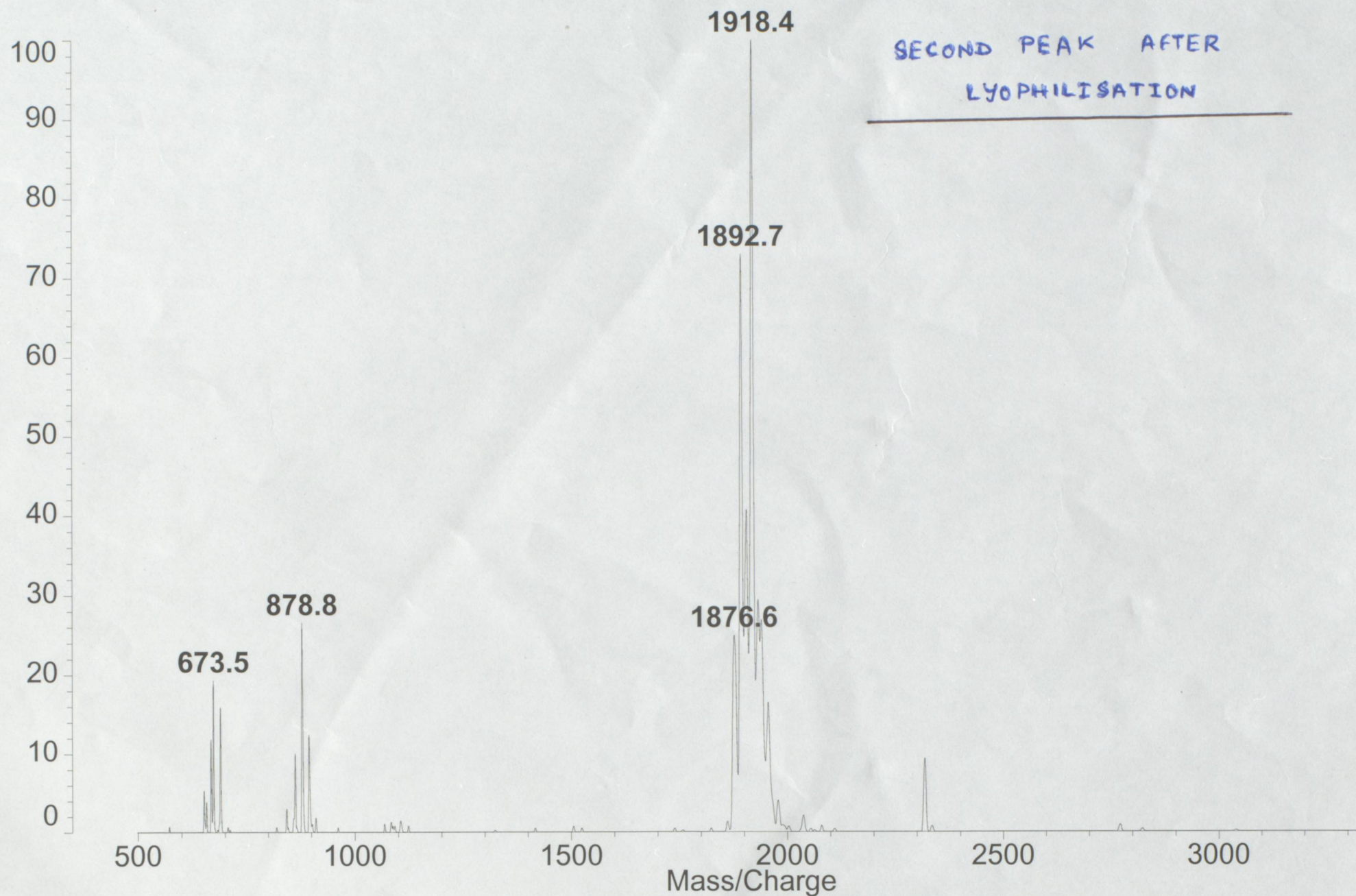


T (min)	CH <sub>3</sub> CN %
0	20
30	40
50	60
70	90
80	95
90	95
100	20

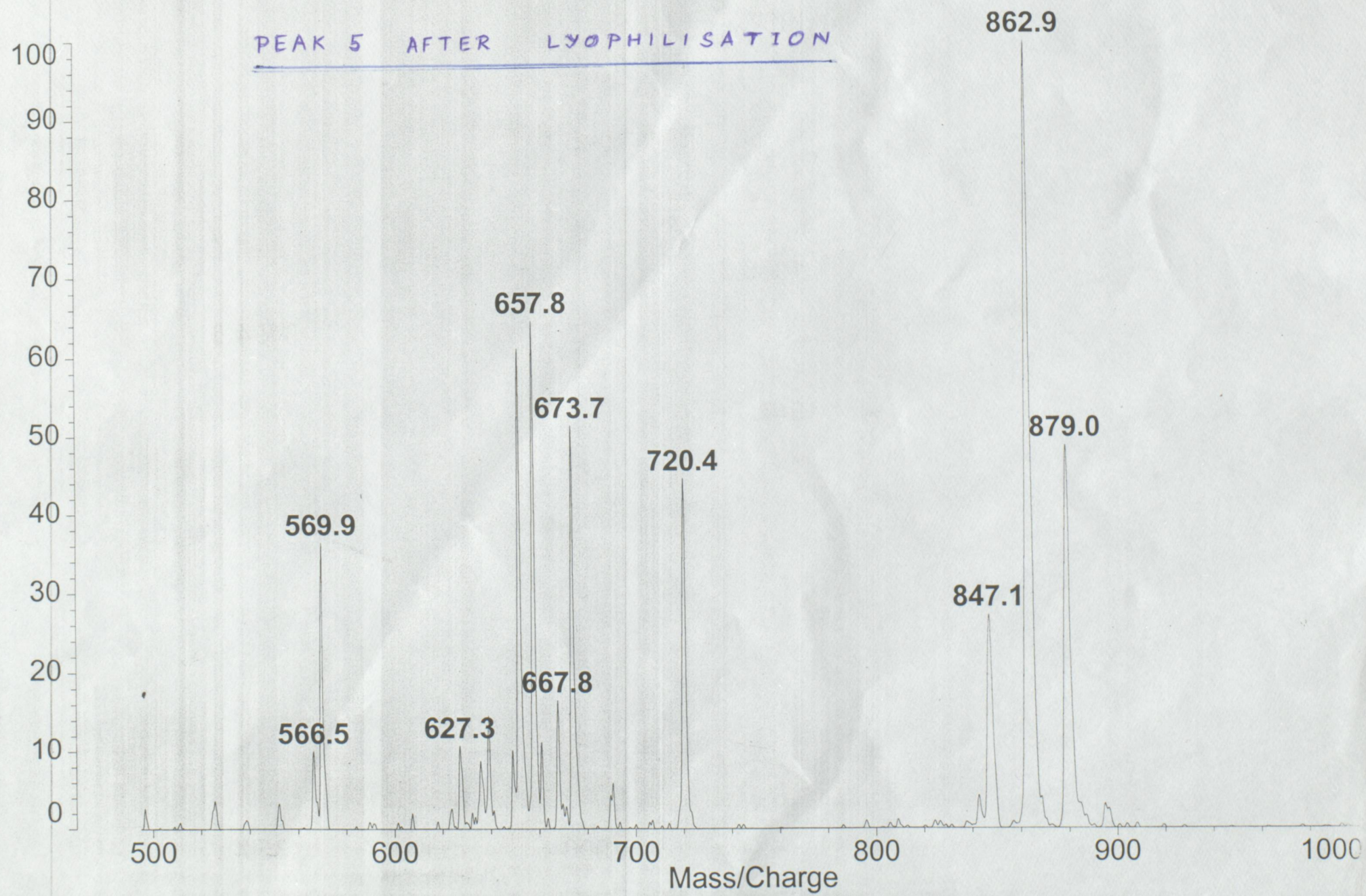
%Int. 100% = 19 mV[sum= 632 mV] Profiles 1-33 Smooth Av 100



%Int. 100% = 5.6 mV[sum= 140 mV] Profiles 1-25 Smooth Av 300 -Baseline 500

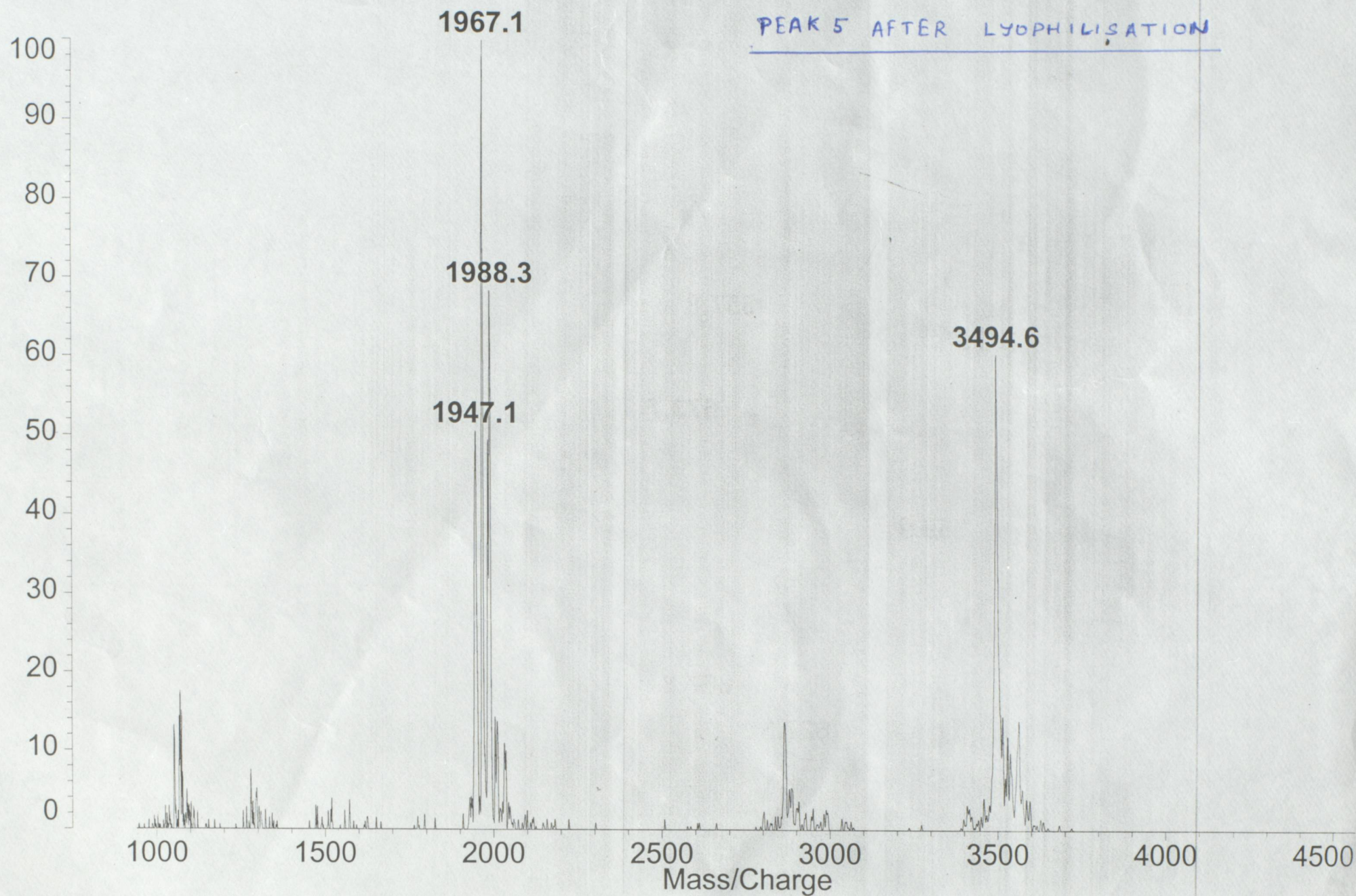


%Int. 100% = 10 mV[sum= 608 mV] Profiles 1-60 Smooth Av 100 -Baseline 100



*5th peak  
after lyophilisation.*

%Int. 100% = 2.3 mV[sum= 136 mV] Profiles 1-60 Smooth Av 100 -Baseline 100



%Int. 100% = 4.8 mV[sum= 290 mV] Profiles 1-60 Smooth Av 50 -Baseline 100

